09/214,752B Does Not Comply Corrected Diskette Needed 61207 delete-totused in new Seguence Rule format L1417 <210> 1 sle sample Sequence Listery <211> 382 <212> PRT (attacked) for valid format <213> Homo sapiens Also, corsult rew Sequence <400> 1 Kules TECH CENTER 1800720 Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

t deral-Kegister / vui. 03. No. 104. Monuay, June 1. 1938 / Kin Scanding Kutauons

29639≥

milt & 10 8 00 par Grof rate Sample Sajur

Smith, Jane:

120> Example of a Sequence Listing

<130> 01-00001

<140> US 08/999,999

<141> 1998-02-28

<150> EP 91000000

<151> 1997-12-31

Please consult,

2. (1605-2)

<170> PatentIn ver. 2.0

<210> 1

<21i> 403

<212> DNA

<213> Paramecium aurelia

<220>

<221> CDS

<222> 341..394

<300>

<301> Doe, Richard

<302> Isolation and Characterization of a Gene Encoding a

Protease from Paramecium sp.

<303> Journal of Fictional Genes

<3.05%

<306> 1 - 7

-<307>#1988-06-20

<400> 1

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ctcttgagtc ctggagatct ctcctctcac atgtgatcgt cgagactgac cgatagatcg 120

ctgactgact ctgagatagt cgagcccgta cgagacccgt cgagggtgac agagagtggg 180

cgcgtgcgcg cagagcgccg cgccggtgcg cgcgcgagtg cgcggtgggc cgcgcgaggg 240

ctttcgcggc agcggcggcg ctttccggcg cgcgcccgtc cgcccctaga cctgagaggt 300

cttctcttcc ctcctcttca ctagagaggt ctatatatac atg gtt tca atg ttc 355

Met Val Ser Met Phe

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age ttg tet tte aaa tgg eet gga ttt tgt ttg ttt gtt tgtttgete

403

Ser Leu Ser Phe Lys Trp Pro Gly Phe Cys Leu Phe Val-

10

<210> 2

<211> 18

<212> PRT

<213>: Paramecium aurelia

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<400> 2

Ser Met Die Serifeurser Phe Lys urd Deo Cly Die Cya Pen

ed: May 22, 1998. · A. Lehman, oc. 98-14194 Filed 5-29-98; 8:45 am] 1 COO€ 3510-16-C

itable. The numeric identifier shall be used only in the Seque.

Listings The order and presentation of the items of information in the "sequence Visting" shall conform to the arrangements given below. Each item of information shall begin on a new line and shall begin With the numeric identifier enclosed in angle brackets as shown. The submission of those items of information designated with an "M" is mandatory. The submission of those items of information designated with an "O" is optional. Numeric identifiers <110> through <170> shall only be set forth at the beginning of the "Sequence Listing." The following table illustrates the numeric identifiers.

pplie tions OG Date 23 June 1998

Numerics designation of the second se	Definition	Comments and Format	Mandatory (M) or Optional (O)
<110>	Applicant	Preferably max. of 10 names; one name per line; preferable format: Surname, Other Names and/or Initials	Н
<120>	Title of Invention	•	м
<130>	File Reference	Personal file reference	M when filed prior to assignment of appl. number
<140>	Current Applica- tion Number	Specify as: US 07/999,999 or PCT/US96/99999	M, if available
<141>	Current Filing Date	Specify as: yyyy-mm-dd	M, if available
<150>	Prior Application Number	Specify as: US 07/999,999 or PCT/US96/99999	M, if applicable include priority documents under 35 USC 119 and 120
<151>	Prior Application Filing Date	Specify as: yyyy-mm-dd	M, if applicable
<160>	Number of SEQ ID NOs	Count includes total number of SEQ ID NOs	M :
<170>	Software	Name of software used to create the Sequence Listing	0
<210>	SEQ ID NO:#:	Response shall be an integer representing the SEQ ID NO shown	М
<211>	Length	Respond with an integer expressing the number of bases or amino acid residues	M →

sequence mole cule 1s DNA RNA Or PRT (ptotein). If a nucleotide sequenc contains_both_DNA and RNA fragments, the type shall be "DNA." In ad-dition, the combined DNA/ RNA molecule shall be further described in the <220> to <223> feature section. М Scientific name, Organism <213> i.e. Genus/species, Unknown or Artificial Sequence. In addition, the "Unknown" or "Artificial Sequence" organisms shall be further described in the <220> to <223> feature section. · M, under the Leave blank after <220> **Feature** following condi-<220>. <221-223> tions: if "n," provide for a "Xaa," or a moddescription of ified or unusual points of bio-L-amino acid or logical signimodified base was ficance in the used in a sesequence. quence; if ORGAN-ISM is "Artificial Sequence" or "Unknown"; if molecule is combined DNA/RNA. M, under the fol-<221> Name/Key Provide appropriate lowing conditions: identifier for if "n," "Xaa," or feature, preferably from a modified or unusual L-amino WIPO Standard ST.25 (1998), acid or modified base was used in Appendix 2, Tables 5 and 6 a sequence M, under the fol-Specify location <222> Location lowing conditions: within sequence; if "n," "Xaa," or . where appropriate a modified or unstate number of first and last usual L-amino

bases/amino acids

acid or modified

Picquirements for Applica	स्टरम ट्रिक्ट क्रिक्स क्रिक्ट क्रिक्ट क्रिक्ट	This is the second of the seco		
		in feature	bose was used for the state of	
<223>	Other-Infor-	Other relevant information; four lines maximum	M;='und'r=the fol- lowing conditions: if 'n, '"Xaa," or a modified or un- usual=L-amino-acid-	
	The second secon		or modified base wassused in a sequence; if the sequence of th	
,	en in the first probabilities	*•	is "Artificial" Sequence" or "Unknown"; if • • molecule is com- bined DNA/RNA.	
<300>	Publication Information	Leave blank after <300>	0 .	
<301>	Authors	Preferably max of ten named authors of publi- cation; specify one name per line;preferable format: Surname, Other Names and/or Initials	RECEIVED NOV 27 2000 TECH CENTER 16000/2000	
<302>	Title		0	
<303>	Journal		0	
<304>	Volume		· •	
<305>	Issue		0	
<306>	Pages		0	
<307>	Date	Journal date on which data published; specify as yyyy-mm-dd, MMM-yyyy or Season-yyyy	0	
<308>	Database Accession Number	Accession number assigned by data-base including database name	0	
<309>	Database Entry Date	Date of entry in database; specify as yyyy-mm-dd or MMM-yyyy	0	
<310>	Patent Document Number	Document number; for patent-type citations only. Specify as, for example, US 07/999,999	o 	

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date forepatent type citations only/.

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j.

<312>

Publication Date Document publicati n

patent-type ecitations only, passpecify as wyyyy mm-dd

<313>

Relevant Residues FROM (posicion) To

<400> Sequence

SEQ ID NO should M
follow the
numeric identifier
and should appear
on the line preceding the actual
sequence

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- 5. Section 1.824 is revised to read as follows:
- 1.824 Form and format for nucleotide and/or amino acid sequence submissions in computer readable form.
- (a) The computer readable form required by 1.821(e) shall meet the following specifications:
- (1) The computer readable form shall contain a single "Sequence Listing" as either a diskette, series of diskettes, or other permissible media outlined in paragraph (c) of this section.
- (2) The "Sequence Listing" in paragraph (a) (1) of this section shall be submitted in American Standard Code for Information Interchange (ASCII) text. No other formats shall be allowed.
- (3) The computer readable form may be created by any means, such as word processors, nucleotide/amino acid sequence editors or other custom computer programs; however, it shall conform to all specifications detailed in this section.
- (4) File compression is acceptable when using diskette media, so long as the compressed file is in a self-extracting format that will decompress on one of the systems described in paragraph (b) of this section.
- (5) Page numbering shall not appear within the computer readable form version of the "Sequence Listing" file.
- (6) All computer readable forms shall have a label permanently affixed thereto on which has been hand-printed or typed: the name of the applicant, the title of the invention, the date on which the data were recorded on the computer readable form, the operating system used, a reference number, and an application serial number and filing date, if known.
- (b) Computer readable form submissions must meet these format requirements:
- (1) Computer: IBM PC/XT/AT, or compatibles, or Apple Macintosh;
- (2) Operating System: MS-DOS, Unix or Macintosh;